

Molecular Identification of *Bartonella* Species in Dogs and Arthropod Vectors in Hamedan and Kermanshah, Iran

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Abstract

Bartonella species are lesser-known pathogenic bacteria that infect a wide range of domestic and wild animals as well as humans. Currently, out of 40 *Bartonella* species/subspecies, at least 17 are associated with clinical signs in humans and animals. However, despite the zoonotic importance of bartonellosis, there is limited information about prevalence and species infecting dogs and cats in Iran. The aim of this study was molecular identification of *Bartonella* species in dogs and arthropods infesting them in Hamedan and Kermanshah cities in the west of Iran. Blood genomic DNA (gDNA) was extracted from 100 dogs (45 from Hamedan and 55 from Kermanshah) and, of 25 *Ctenophalides canis* fleas, six *Pulex irritans* fleas and 12 *Rhipicephalus sanguineus* ticks collected from nine infested dogs were examined for the presence of *Bartonella* species. Conventional PCR targeting fragments of ITS and *rpoB* genes was performed, and PCR-positive samples were sequenced bidirectionally and analyzed phylogenetically. Out of 100 dogs, 14 dogs (14%, ten from Hamedan and four from Kermanshah) were found infected with *Bartonella* species. Nucleotide sequencing confirmed the presence of four *Bartonella* species in the examined population i.e. *Bartonella vinsonii* subsp. *berkhoffii* and *Candidatus B. merieuxii* in dogs. None of the examined fleas scored positive but one *Rh. sanguineus* tick from a blood-negative dog was infected with *Bartonella* DNA. Results of the present study showed the presence of different zoonotic *Bartonella* species in dogs of Hamedan and Kermanshah cities highlighting the importance of this vector-borne infection. Effective ectoparasite control strategies, regular examination of pet and urban dogs and cats and successful chemoprophylaxis are suggested.

Key words: Arthropods, *Bartonella*, Dogs, Hamedan, Kermanshah, PCR, Zoonotic

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